

<b>Notice of References Cited</b>	Application/Control No. 10/091,019	Applicant(s)/Patent Under Reexamination HARRIS ET AL.	
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**U.S. PATENT DOCUMENTS**

*		Document Number Country Code-Number-Kind Code	Date MM-YYYY	Name	Classification
*	A	US-6,790,443	09-2004	Stern et al.	424/130.1
*	B	US-6,555,346	04-2003	Kretzdorn et al.	435/71.1
	C	US-			
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	J	US-			
	K	US-			
	L	US-			
	M	US-			

**FOREIGN PATENT DOCUMENTS**

*		Document Number Country Code-Number-Kind Code	Date MM-YYYY	Country	Name	Classification
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**NON-PATENT DOCUMENTS**

*		Include as applicable: Author, Title Date, Publisher, Edition or Volume, Pertinent Pages)
	U	Taticek et al, 1997. Biotechnol. Bioeng. 54: 142-152. ✓
	V	Wells (18 September 1990) "Additivity of Mutational Effects in Proteins." Biochemistry 29(37): 8509-8517. ✓
	W	Ngo et al. (2 March 1995) "The Protein Folding Problem and Tertiary Structure Prediction, Chapter 14: Computational Complexity Protein Structure Prediction, and the Levinthal Paradox" pp. 492-495. ✓
	X	Bork (2000) "Powers and Pitfalls in Sequence Analysis: The 70% Hurdle." Genome Research 10:398-400. ✓

\*A copy of this reference is not being furnished with this Office action. (See MPEP § 707.05(a).)  
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	V	Doerks et al. (June 1998) "Protein annotation: detective work for function prediction." Trends in Genetics 14(6): 248-250. ✓
	W	Smith and Zhang (November 1997) "The challenges of genome sequence annotation or 'The devil is in the details'." Nature Biotechnology 15:1222-1223. ✓
	X	Brenner (April 1999) "Errors in genome annotation." Trends in Genetics 15(4): 132-133. ✓

\*A copy of this reference is not being furnished with this Office action. (See MPEP § 707.05(a).)  
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